



Replacement Sheet

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1 GTCCCTCCACCATGCACTCGCTGGGCTTCTTCTGTGGCGTGTCTCTCTGCTCGCCGCTG 60
-----+-----+-----+-----+-----+-----+-----+-----+
CAGGAAGGTGGTACGTAGCGACCCGGAAGAAGAGACACCCGACCAAGAGACGAGCGCGGAC
M H S L G F F S V A C S L L A A A -
CGCTGCTCCGGGTCTCGGAGGCGCCGCGCGCGCGCGCTTTCGAGTCCGGACTCG 120
-----+-----+-----+-----+-----+-----+-----+-----+
GCGACGAGGCGCCAGGAGCGCTCCGCGGCGCGCGCGGCGGAGCTCAGGCCCTGAGC
L L P G P R E A P A A A A A F E S G L D -
ACCTCTCGGACGCGGAGCCCGACGCGGCGGAGGCCACGGCTTATGCAAGCAAAGATCTGG 180
-----+-----+-----+-----+-----+-----+-----+-----+
TGGAGAGCCCTGCGCCTCGGCGTGGCGCCCGCTCCGGTGCCGAATACGTTCGTTCTAGACC
L S D A E P D A G E A T A Y A S K D L E -
AGGAGCAGTTACGGTCTGTGTCCAGTGTAGATGAACCTCATGACTGTACTCTACCCAGAAT 240
-----+-----+-----+-----+-----+-----+-----+-----+
TCCTCGTCAATGCCAGACACAGGTCACATCTACTTGAGTACTGACATGAGATGGGTCTTA
E Q L R S V S S V D E L M T V L Y P E Y -
ATTGGAATAATGTACAAGTGTGCTAGCTAAGGAAAGGAGGCTGGCAACATAACAGAGAACAGG 300
-----+-----+-----+-----+-----+-----+-----+-----+
TAACCTTTACATGTTACAGTCGATTCCTTTCCTCCGACCGTGTGTATTTGTCTCTGTCTCC
W K M Y K C Q L R K G G W Q H N R E Q A -
CCAACCTCAACTCAAGGACAGAGAGACTATATAAATTGCTGCAGCACATTATAATACAG

FIG. 1A



Replacement Sheet

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TTTACAGACAAGTTTCATTCCATTATTAGACGTTCCCTGCCAGCAACACTACCACAGTGTC 720
-----+-----+-----+-----+-----+-----+-----+
AAATGCTCTGTTCAAGTAAGGTAATAATCTGCAAGGACGGTCGTTGTGATGGTGTACACAG
Y R Q V H S I I R R S L P A T L P Q C Q -
AGGCAGCGAACAAGACCTGCCCCACCAATTACATGTGGAATAATCACATCTGCAGATGCC 780
-----+-----+-----+-----+-----+-----+-----+
TCCGTCGCTTGTCTGGACGGGGTGGTTAATGTACACCTTATTAGTGTAGACGCTCTACGG
A A N K T C P T N Y M W N N H I C R C L -
TGGCTCAGGAAGATTTTATGTTTTCCTCGGATGCTGGAGATGACTCAACAGATGGATTCC 840
-----+-----+-----+-----+-----+-----+-----+
ACCGAGTCCTTCTAAATAACAAAGGAGCCTACGACCTCTACTGAGTTGTCTACCTAAGG
A Q E D F M F S S D A G D S T D G F H -
ATGACATCTGTGGACCAACAAGAGCTGGATGAAGAGACCTGTCAGTGTGTCTGCAGAG 900
-----+-----+-----+-----+-----+-----+-----+
TACTGTAGACACCTGGTTTGTTCCTCGACCTACTTCTCTGGACAGTCAACACAGACGTCTC
D I C G P N K E L D E E T C Q C V C R A -
CGGGGCTTCGGCCTGCCAGCTGTGGACCCCAAGAACTAGACAGAAACTCATGCCAGT 960
-----+-----+-----+-----+-----+-----+-----+
GCCCCGAAGCCGACGGTCGACACCTGGGGTGTCTCTTGATCTGTCTTTGAGTACGGTCA
G L R P A S C G P H K E L D R N S C Q C -
GTGTCTGTAAAAACAACACTCTTCCCCAGCCCAATGTGGGGCCCAACCGAGAAATTGATGAAA 1020
-----+-----+-----+-----+-----+-----+-----+
CACAGACATTTTGTGTTGAGAAGGGGTCGGTTACACCCCGGTGGCTCTTAAACTACTTT

FIG. 1C



V C K N K L F P S Q C G A N R E F D E N -
ACACATGCCAGTGTGTATGTAAGAACCTGCCCCAGAAATCAACCCCTAAATCCTGGAA 1080
-----+-----+-----+-----+-----+-----+-----+
TGTGTACGGTCACACATACATTTCTTGGACGGGCTCTTTAGTTGGGATTTAGGACCTT
T C C Q C V C K R T C P R N Q P L N P G K -
AATGTGCCCTGTGAATGTACAGAAAGTCCACAGAAATGCTTGTAAAGGAAAGAGTTCC 1140
-----+-----+-----+-----+-----+-----+-----+
TTACACGGACACTTACATGTCTTTTCAGGTGTCTTTACGAACAATTTTCCCTTTCTTCAAGG
C A C E C T E S P Q K C L L K G K K F H -
ACCACCAACATGCAGCTGTACAGACGGCCCATGTACGAACCGCCAGAAAGGCTTGTGAGC 1200
-----+-----+-----+-----+-----+-----+-----+
TGGTGGTTTGTACGTCGACAAATGTCTGCCGGTACATGCTTGGCGGTCTTCCGAAACACTCG
H Q T C S C Y R R P C T N R Q K A C E P -
CAGGATTTTCATATAGTGAAGAAGTGTGTCGTTGTGTCCCTTCATATTGGCAAAGACCAC 1260
-----+-----+-----+-----+-----+-----+-----+
GTCCCTAAAAGTATATCACTTCTTCACACAGCAACACAGGGAAGTATAACCGTTTCTGTG
G F S Y S E E V C R C V P S Y W Q R P Q -
AAATGAGCTAAGATTGTACTGTTTCCAGTTCATCGATTTTCTATTATGGAAAACTGTGT

FIG. 1D



Replacement Sheet

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1261 -----+-----+-----+-----+-----+-----+-----+ 1320
TTTACTCGATTCTAACATGACACAAAAGGTCAAGTAGCTAAAGATAATACCTTTTGGACACA
M S *
1321 TGCCACAGTAGAACTGTCTGTGAACAGAGAGACCCCTTGTGGTGCCATGCTAACAAAGACA 1380
ACGGTGTCATCTTGACAGACACTTGTCTCTCTGGGAACACCCAGGTACGATTGTTTCTGT
AAAGTCTGTCTTTCCCTGAACCATGTGGATAACTTTACAGAAATGGACTGGAGCTCATCTG
1381 -----+-----+-----+-----+-----+-----+-----+ 1440
TTTCAGACAGAAAGGACTTGGTACACCTATTGAAATGTCTTTACCTGACCTCGAGTAGAC
CAAAAGGCCCTCTTGTAAGACTGGTTTCTGCCAATGACCAACAGCCAAGATTTTCCTC
1441 -----+-----+-----+-----+-----+-----+-----+ 1500
GTTTCCGGAGAACATTCTCGACCACAAAGACGGTTACTGGTTGTCTGCTTCTAAAAGGAG
TTGTGATTCTTTAAAGAATGACTATATAATTTATTTCCACTAAAAATATTGTTTCTGTC
1501 -----+-----+-----+-----+-----+-----+-----+ 1560
AACACTAAAGAAATTTTCTTACTGATATATAATAAAGGTGATTTTATAACAAAGACG
ATTCATTTTATAGCAACAATAATTGGTAAACTCACTGTGATCAATAATTTTATATCAT
1561 -----+-----+-----+-----+-----+-----+-----+ 1620
TAAGTAAATAATCGTTGTTGTTAACCATTTTGAGTGACACTAGTTATAAAAATATAGTA
GCAAAAATATGTTTAAATAAAATGAAATTTGTATTTATAAAAAAAA
1621 -----+-----+-----+-----+-----+-----+-----+ 1674
CGTTTATACAAATTTTATTTTACTTTTACATAAATAATTTTCTTTTCTTTT

FIG.1E